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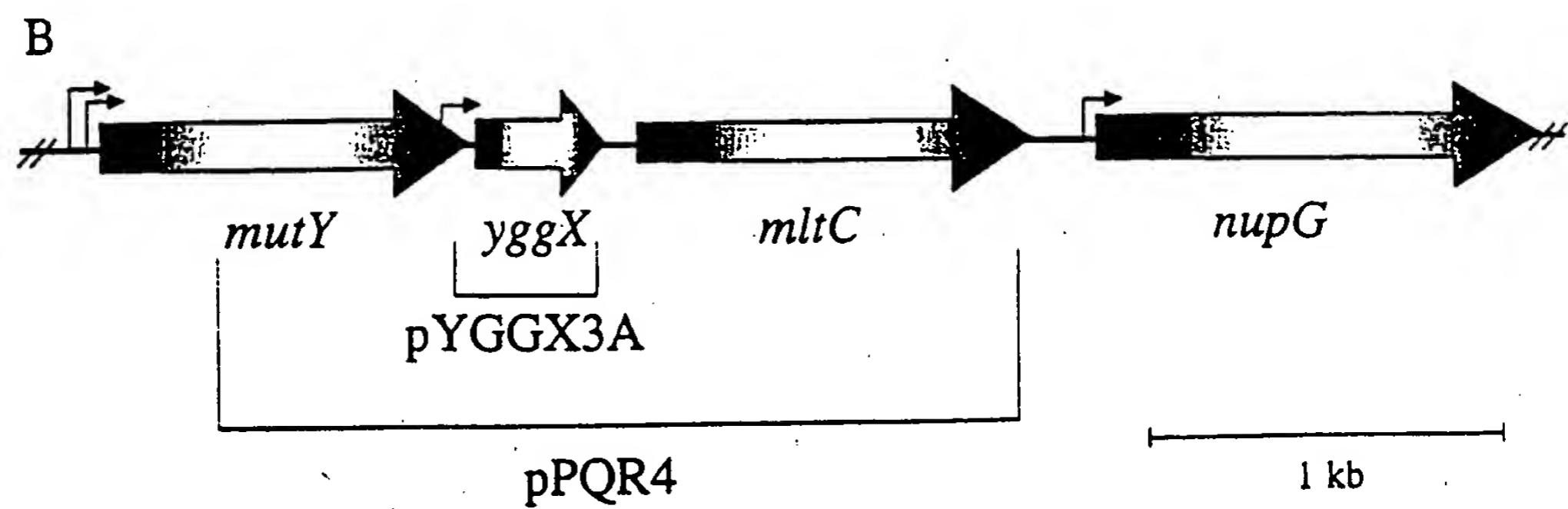


Fig. 1. Physical parameters of *yggX* and its gene product. (A) Alignment of *YggX* homologs. (B) Operon structure of *mutY/yggX* in *E. coli* and *S. enterica* LT2. Promoters were mapped by Gifford and Wallace in *E. coli* (43).

Bpertussis
Bparapert
Bbronchi
A.actin
Pmultocida
Hinfluenzae
Hducreyi
Sputrefasciens
Vcholerae
Ecoli
O157_H7EDL933
O157_H7
Spara
Senteritidis
Sdublin
Styphict18
Styphimurium
Kpneumo
Ypesits
Buchnera
Xfastidiosa
Psyring
Pputida
Paeruginosa
Ngonorrhoeae
NmeningitB
NmeningitA
Bmallei
Bpseudomallei
Tferrooxidans
Mcapsulatus
Cburnetii

1 MSRIVNCVKLKREAEGLDFPPYPGE LGTRIWQQI SKEAWEWKQIQTRLVNEENRLNLADA
 1 MSRIVNCVKLKREAEGLDFPPYPGE LGTRIWQQI SKEAWEWKQIQTRLVNEENRLNLADA
 1 MSRIVNCVKLKREAEGLDFPPYPGE LGTRIWQQI SKEAWEWKQIQTRLVNEENRLNLADA
 1 MARMVFCERLKQEAEGLDFQLYPGE LGKRIEFDISKQAWGEWMKKQTMLVNEKKLNMMNA
 1 MARTVFCEYLKQESEGLDFQLYPGE LGKRIEFDISKQAWREWMKKQTMLVNEKKLNMMNA
 1 MARTVFCEYLKKEAEGLDFQLYPGE LGKRIEFDISKQAWGEWIKKQTMLVNEKKLNMMNA
 1 MARMVFCYLNKEADGLDFQLYPGDLGKRIEFDISKQAWAEWIKKQTMLVNEKKLNMMNP
 1 MARTVNCVHLNKEADGLDFQLYPGDLGKRIEFDISKQAWGLWQKKQTMLVNEKKLNMMNV
 1 MARTVFCTRLQKEADGLDFQLYPGE LGKRIEFDISKQAWAQAQWQHKQTMLVNEKKLNMMDP
 1 MSRTIFCTFLQREAEGQDFQLYPGE LGKRIYNEISKQAWAQAQWQHKQTMLVNEKKLNMMNA
 1 MSRTIFCTFLQREAEGQDFQLYPGE LGKRIYNEISKQAWAQAQWQHKQTMLVNEKKLNMMNA
 1 MSRTIFCTFLQREAEGQDFQLYPGE LGKRIYNEISKQAWAQAQWQHKQTMLVNEKKLNMMNA
 1 MSRTIFCTYLQRDAEGQDFQLYPGE LGKRIYNEISKDAWAQAQWQHKQTMLVNEKKLNMMNA
 1 MSRTIFCTYLQRDAEGQDFQLYPGE LGKRIYNEISKDAWAQAQWQHKQTMLVNEKKLNMMNA
 1 MSRTIFCTYLQRDAEGQDFQLYPGE LGKRIYNEISKDAWAQAQWQHKQTMLVNEKKLNMMNA
 1 MSRTIFCTYLQRDAEGQDFQLYPGE LGKRIYNEISKDAWAQAQWQHKQTMLVNEKKLNMMNA
 1 MSRTIFCTFLQREADGQDFQLYPGE LGKRIYNEISKQAWAQAQWQHKQTMLVNEKKLSMMNP
 1 MSRTIFCTFLKKDAERQDFQLYPGEIGKRIYNEISKQAWSQWITKQTMLVNEKKLSMMNI
 1 MNRIIFCTFFKKSEGQDFQSYPGKLGKKIYDQISKKAWEKWIEKQTILINEENLNFNL
 1 MORIIIFCEYEQRDTEGLDFVPYPGE LGQKIFACIGKVGWAAWLVHQTMLVNEENRLSPRNP
 1 MTRTVMCRKYKEELPGLERAPYPGAKGEDIFNHVSQKA WADWQKHQTLLVNEERRLNMMNA
 1 MTRTVMCRKYQEELPGLERPPYPGAKGQDIFEHISKQAWADWQKHQTMLVNEKRLNMMNA
 1 MSRTVMCRKYHEELPGLDRPPYPGAKGEDIYNNVSRKAWDEWQKHQTMLVNEERRLNMMNA
 1 MARMVFCVKLKREAEGMKFPPLPNELGKRIEFDENVSQEAWAAWTRHQTMLVNEENRLSLADP
 1 MARMVFCVKLKREAEGMKFPPLPNELGKRIEFDENVSQEAWAAWTRHQTMLVNEENRLSLADP
 1 MARMVFCVKLKREAEGMKFPPLPNELGKRIEFDENVSQEAWAAWTRHQTMLVNEENRLSLADP
 1 MARMIHCAKLGKEAEGLDFPPLPGE LGKRLYESVSKQAWQDWLKQQTMLVNEENRLNMADP
 1 MARMIHCAKLGKEAEGLDFPPLPGE LGKRLYESVSKQAWQDWLKQQTMLVNEENRLNMADP
 1 MSRMVQCVKLGHEAEGLDRPPYPGALGARIYQEVSKEAWQGWLKHQTMLVNEYRLSPIDP
 1 MARRIIICAKLGIEADGLDAPPFPGPQGQRIFEEHVSKEAWQDWLKLQQTMLVNEHRLTPFEA
 1 MTRRIICQKLGKEADALNYSPYPGELGERIYNHISEQAWQAWLSHQTMLVNEYRLSLIDP

Fig. 1A

10
20
30
40
50
60
70
80
90

Bpertussis	61 RARKYLQQQMERELFEDGTVEAQGYVP----
Bparapert	61 RARKYLQQQMERELFEDGTVEAQGYVP----
Bbronchi	61 RARKYLQQQMERELFEDGTVEAQGYVP----
A.actin	61 EHRKLLQEMVNELFEGKDVHIEGYTPPEAK
Pmultocida	61 DHRQLLEQEMVNELFEGKDVHIEGYVP----
Hinfluenzae	61 EHRKLLQEMVNELFEGKDVHIEGYVP----
Hducreyi	61 EHRQLLEAEMVNELFEGKDVHIDGYVP----
Sputrefasciens	61 DDRKFLEAQMTSELFEGKDVEIEGFVPE---
Vcholerae	61 EHRKLLQEMVNELFEGKEVHIEGYTPPAK-
Ecoli	61 EHRKLLQEMVNELFEGKEVHIEGYTPEDKK
O157_H7EDL933	61 EHRKLLQEMVNELFEGKEVHIEGYTPEDKK
O157_H7	61 EHRKLLQEMVNELFEGKEVHIEGYTPEDKK
Spara	61 EHRKLLQEMVNELFEGKEVHIEGYTPEDKK
Senteritidis	61 EHRKLLQEMVSELFEGKDVEIEGYTPEDKK
Sdublin	61 EHRKLLQEMVSELFEGKDVEIEGYTPEDKK
Styphict18	61 EHRKLLQEMVSELFEGKDVEIEGYTPEDKK
Styphimurium	61 EHRKLLQEMVSELFEGKDVEIEGYTPEDKK
Kpneumo	61 EHRKLLQEMVQELFEGK-----
Ypesits	61 EDRKLLEQEMVNELFEGQDVHIAGYTPPSK-
Buchnera	61 EHRKKIEKYMKLFLK-----
Xfastidiosa	61 SHRAFLEEELNKEFERRVAKPEGYIEPD--
Psyring	61 EDRKFLQTEMKELSGEEYAAQAEQYVPPEK-
Pputida	61 EDRKFLQAEQMDKEFAGEEYAAQAEQYVP---
Paeruginosa	61 EDRKFLQQEMDKELSGEDYAKADGYVP---
Ngonorrhoeae	61 RAREYLAQQMEQYFFGDGADAVQGYVPQ---
NmeningitB	61 RAREYLAQQMEQYFFGDGADAVQGYVPQ---
NmeningitA	61 RAREYLAQQMEQYFFGDGADAVQGYVPQ---
Bmallei	61 RARQYLMKQTEKYFFGEGADQASGYVP---
Bpseudomallei	61 RARQYLMKQTEKYFFGEGADQASGYVP---
Tferrooxidans	61 KSRTFLEKQMEAYFFGDGAQSPEGYVP----
Mcapsulatus	61 SARKFLEQEREKELFGGGTSTPQGYVP----
Cburnetii	61 KARQFLEQEMINELFGTGSEKPGYTSE---

Fig. 1A (continued)

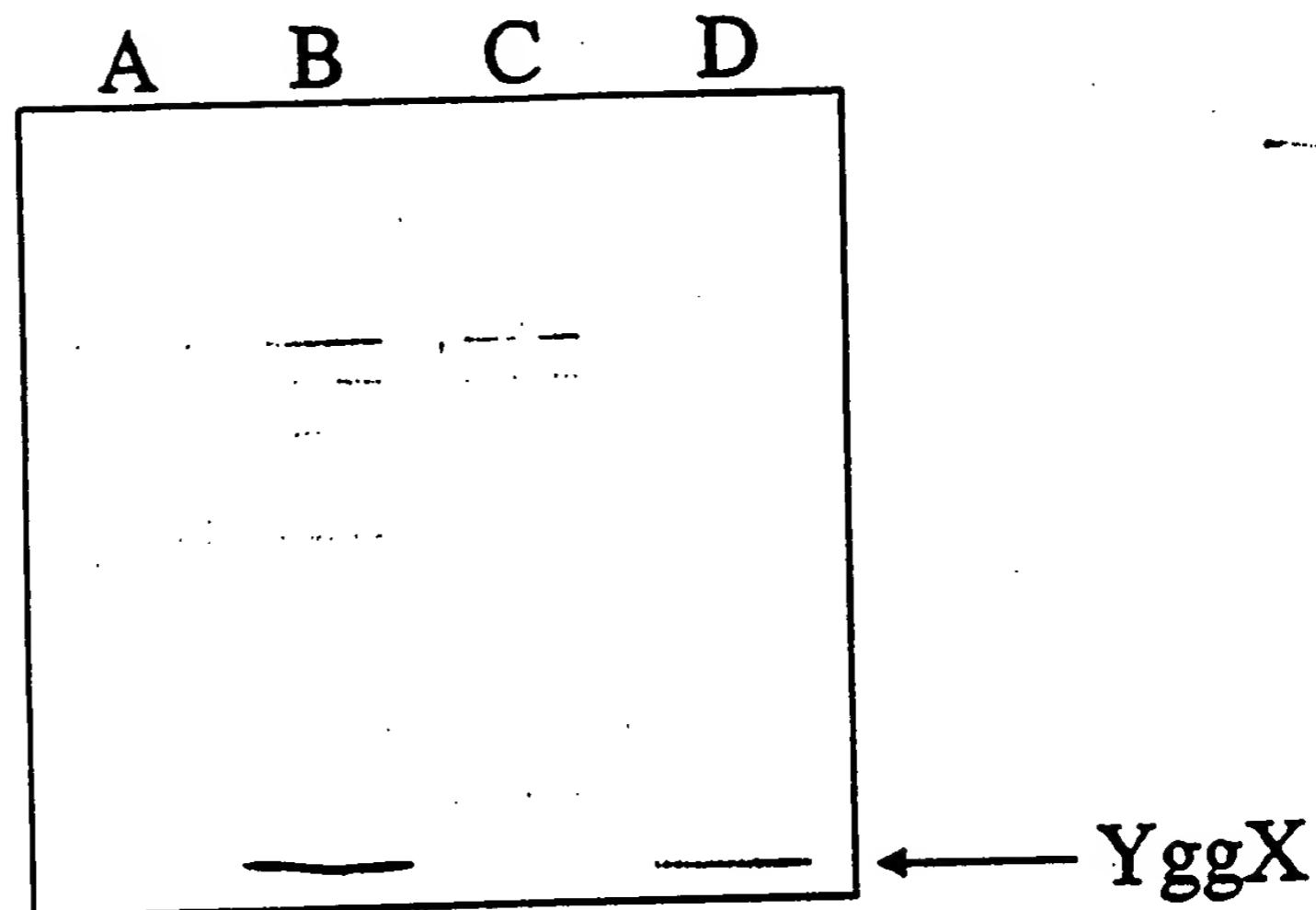


Fig. 2. Increased levels of YggX protein in *yggX** mutant. Western blot analysis was performed according to Harlow and Lane (59). Proteins were visualized by using alkaline phosphatase conjugated to anti-rabbit secondary antibody (Promega). Lanes A-C were loaded with crude cell-free extracts (1 μ g protein) from strains DM5104, DM5105 (*yggX**), and DM5647 (*yggX*::Gm), respectively. Lane D was loaded with 1 ng purified YggX.

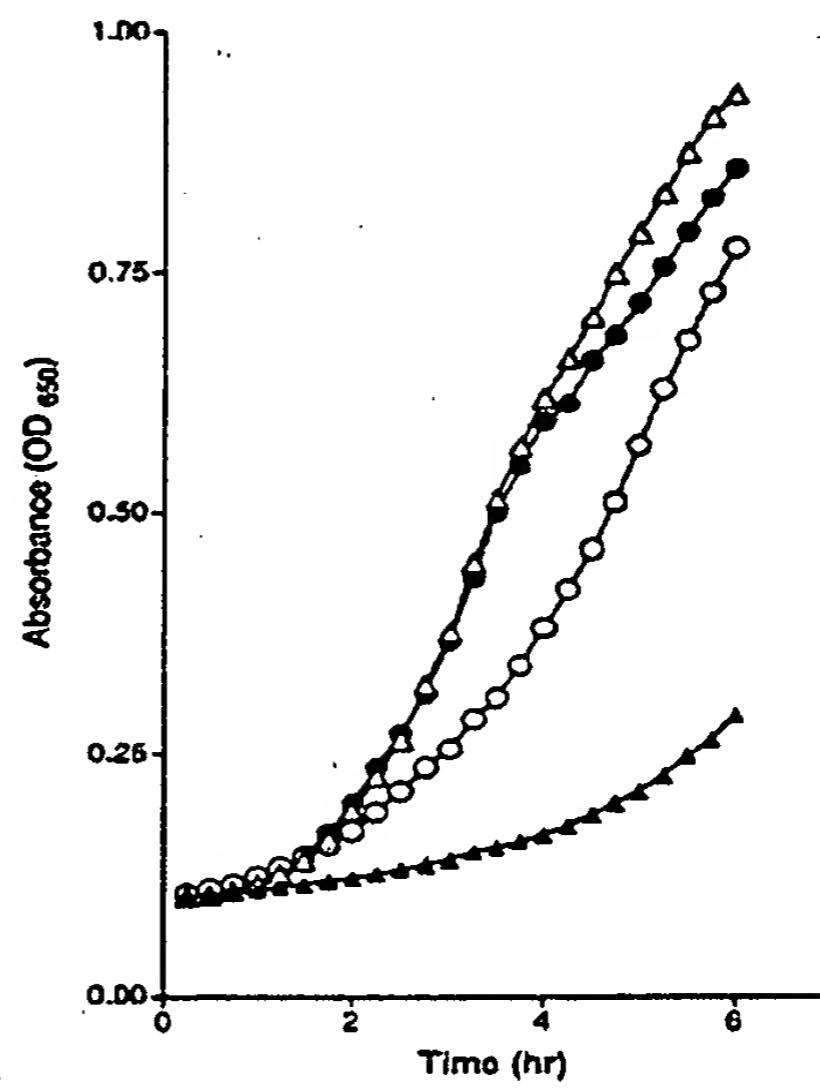


Fig. 3. The *yggX** mutation does not increase MNNG resistance of *gshA* mutants. Strain LT2 was grown in LB with (\blacktriangle) and without (\triangle) 60 μ M MNNG. Both *gshA* (\circ) and *gshA yggX** (\bullet) mutant strains were grown in LB with 60 μ M MNNG.

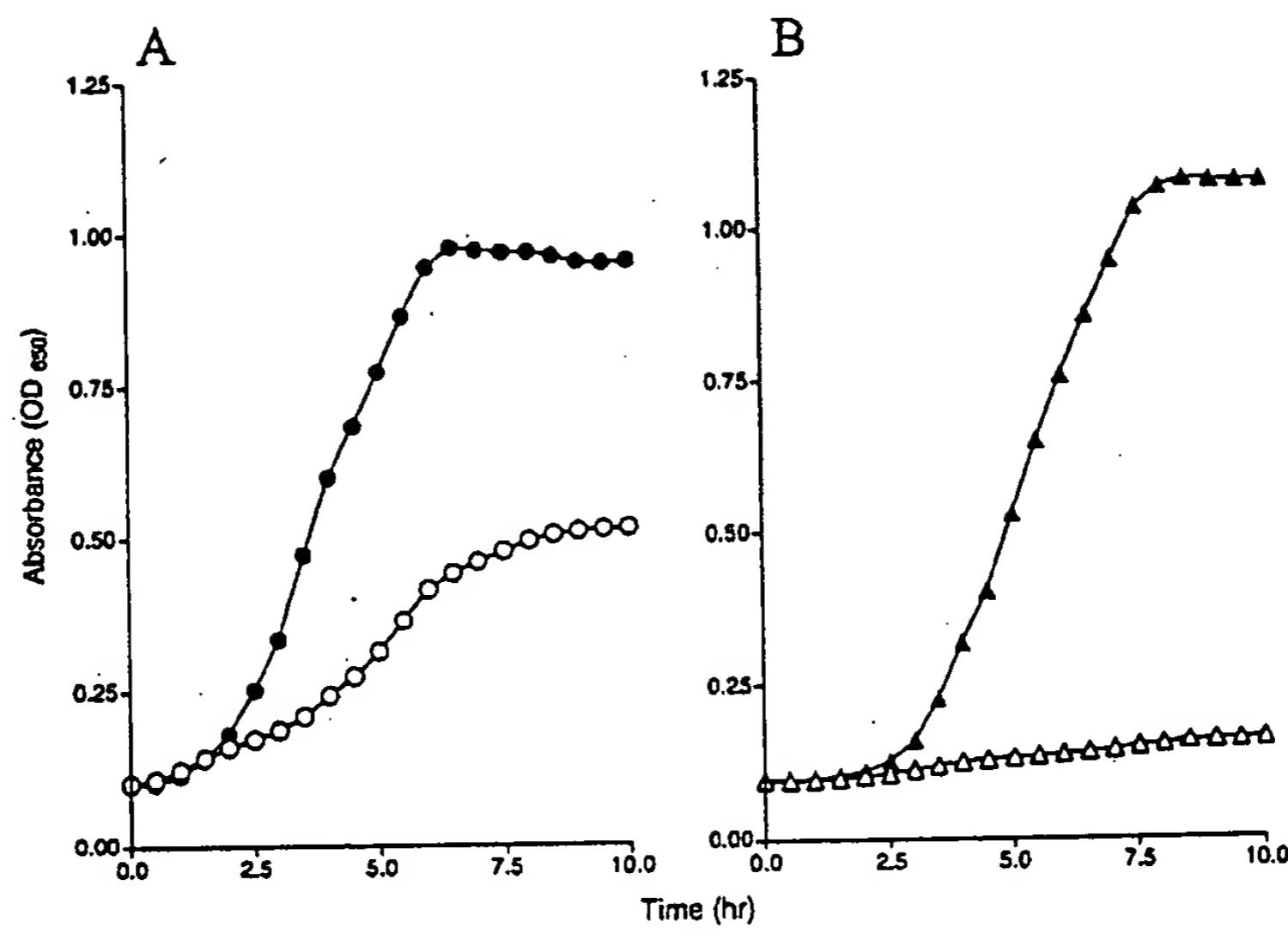


Fig. 4. The *yggX** mutation increases resistance of *S. enterica* to PQ. (A) Growth of *gshA* (○) and *gshA yggX** (●) mutant strains in LB with 4 μ M PQ. (B) Growth of LT2 (△) and *yggX** (▲) strains in LB with 40 μ M PQ.

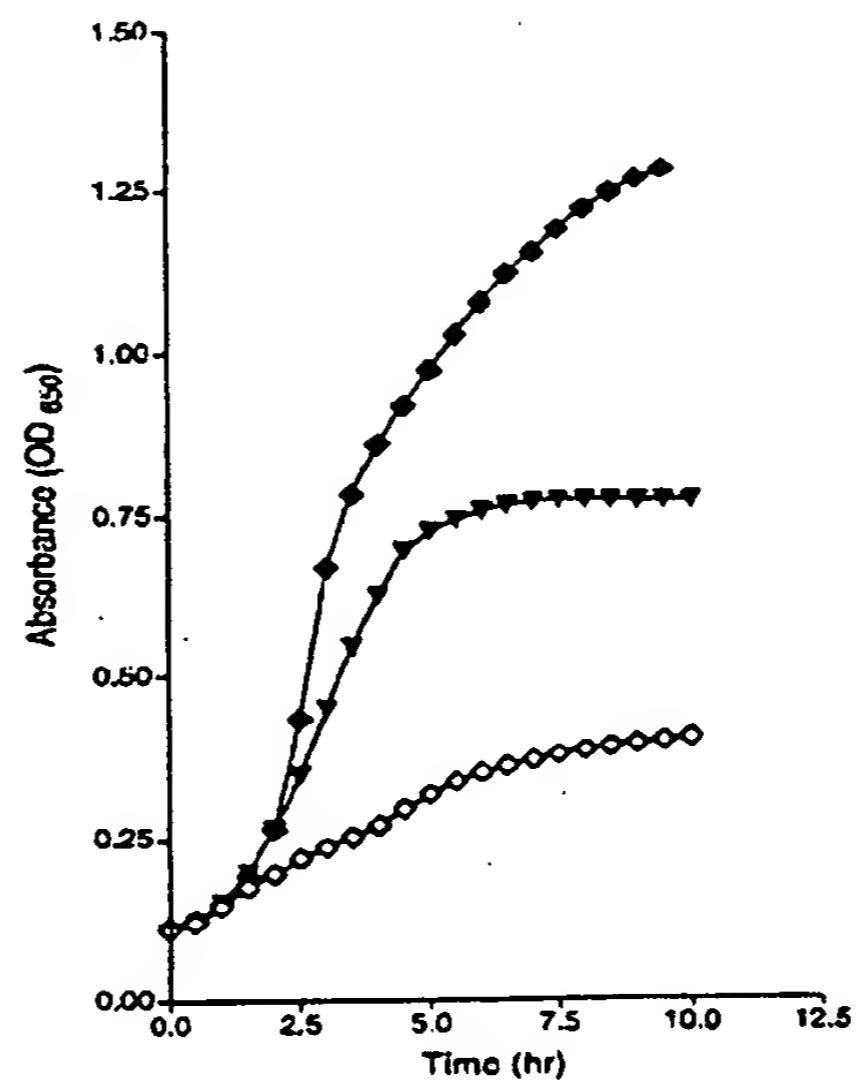


Fig. 5. *yggX** does not require *soxR* to mediate resistance to PQ. Strains LT2 (◆), *soxR* (○), and *soxR yggX** (▼) were grown in LB with 4.0 μ M PQ.

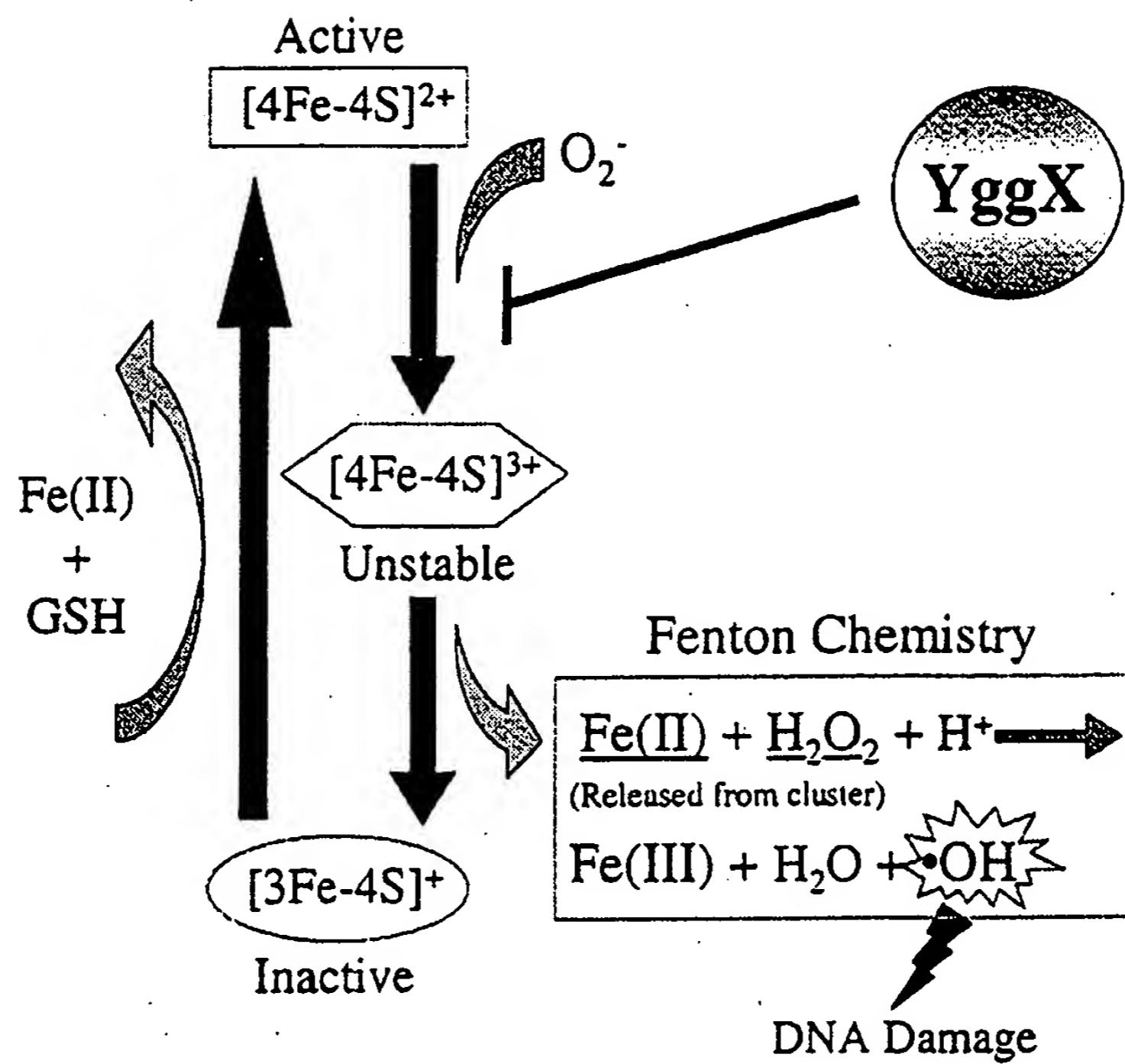


Fig. 6. Model showing how YggX protects *S. enterica* from oxidative damage. The result of superoxide attack on [Fe-S] clusters is depicted. We hypothesize that YggX is able to block oxidative damage to labile clusters and thus prevent the normal downstream consequences of such oxidation.